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- ☐ 1. Document ID: US 5948682 A
Entry 1 of 4 File: USPT Sep 7, 1999
US-PAT-NO: 5948682
DOCUMENT-IDENTIFIER: US 5948682 A
TITLE: Preparation of heterologous proteins on oil bodies

| Full | Title | Citation | Front | Review | Classification | Date | Reference | Claims | KWIC | Image |
|------|-------|----------|-------|--------|----------------|------|-----------|--------|------|-------|
|------|-------|----------|-------|--------|----------------|------|-----------|--------|------|-------|

- ☐ 2. Document ID: US 5830448 A
Entry 2 of 4 File: USPT Nov 3, 1998
US-PAT-NO: 5830448
DOCUMENT-IDENTIFIER: US 5830448 A
TITLE: Compositions and methods for the treatment of tumors

| Full | Title | Citation | Front | Review | Classification | Date | Reference | Claims | KWIC | Image |
|------|-------|----------|-------|--------|----------------|------|-----------|--------|------|-------|
|------|-------|----------|-------|--------|----------------|------|-----------|--------|------|-------|

- ☐ 3. Document ID: US 5843705 A
Entry 3 of 4 File: USPT Dec 1, 1998
US-PAT-NO: 5843705
DOCUMENT-IDENTIFIER: US 5843705 A
TITLE: Transgenically produced antithrombin III

| Full | Title | Citation | Front | Review | Classification | Date | Reference | Claims | KWIC | Image |
|------|-------|----------|-------|--------|----------------|------|-----------|--------|------|-------|
|------|-------|----------|-------|--------|----------------|------|-----------|--------|------|-------|

- ☐ 4. Document ID: US 5762921 A
Entry 4 of 4 File: USPT Jun 9, 1998
US-PAT-NO: 5762921
DOCUMENT-IDENTIFIER: US 5762921 A
TITLE: Composition and methods for the treatment of tumors

| Full | Title | Citation | Front | Review | Classification | Date | Reference | Claims | KWIC | Image |
|------|-------|----------|-------|--------|----------------|------|-----------|--------|------|-------|
|------|-------|----------|-------|--------|----------------|------|-----------|--------|------|-------|

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Content Information for 09/143155

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| Num | Type | Date | Code | Contents Description |
|-----|------|------------|------|--|
| 7 | D | 12/13/1999 | DOCK | DATE CASE WAS DOCKETED |
| 6 | I | 12/02/1998 | M844 | PRIOR ART CITATION FILED P/E |
| 5 | D | 10/24/1998 | DOCK | DATE CASE WAS DOCKETED |
| 4 | I | 08/28/1998 | A.PE | PRE-EXAMINATION AMENDMENT |
| 3 | I | 09/22/1998 | OIPE | APPLICATION DISPATCHED FROM PRE-EXAM |
| 2 | E | 09/08/1998 | SCAN | APPLICATION SCANNED AND DISPATCHED FROM PRE-EXAM |
| 1 | E | 09/03/1998 | IEXX | INITIAL EXAM TEAM XX |

Serial Info

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12/13

L3 ANSWER 1 OF 24 CAPLUS COPYRIGHT 1999 ACS
 AN 1999:656017 CAPLUS
 DN 131:282377
 TI Engineering protein posttranslational modification in **transgenic**
 non-human mammals
 IN Lubon, Henryk; Drohan, William N.; Paleyanda, Rekha K.
 PA American Red Cross, USA
 SO U.S., 20 pp., Cont.-in-part of U.S. 5,589,604.
 CODEN: USXXAM
 DT Patent
 LA English
 IC ICM A01K067-00
 ICS A01K067-027; C12P021-04; C12P021-06
 NCL 800014000
 CC 3-1 (Biochemical Genetics)
 FAN.CNT 8

| | PATENT NO. | KIND | DATE | APPLICATION NO. | DATE |
|----|---|------|----------|-----------------|----------|
| PI | US 5965789 | A | 19991012 | US 1995-434834 | 19950504 |
| | US 5831141 | A | 19981103 | US 1992-943246 | 19920910 |
| | US 5589604 | A | 19961231 | US 1994-247484 | 19940523 |
| | WO 9634966 | A2 | 19961107 | WO 1996-US6121 | 19960506 |
| | W: AU, CA, JP, MX | | | | |
| | RW: AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, | | | | |

SE
 CA 2220109 AA 19961107 CA 1996-2220109 19960506
 AU 9663474 A1 19961121 AU 1996-63474 19960506

PRAI US 1991-638995 19910111
 US 1992-943246 19920910
 US 1994-198068 19940208
 US 1994-247484 19940523
 US 1995-434834 19950504
 WO 1996-US6121 19960506

AB The invention relates to **transgenic** non-human multicellular organisms that contain polynucleotides for expressing proteins that alter posttranslational modification. In particular, the invention provides multiply-**transgenic** animals in which a first **transgene** encodes a first protein, a second **transgene** encodes a second protein, and expression of the second protein affects the posttranslational modification of the first protein in cells of said organism. Expression in preferred embodiments is in specific cells and the modified protein is secreted into a bodily fluid. An example provides

transgenic mice which produce human protein C and the processing protease PACE/furin in mammary glands and secrete both proteins into milk.

The protein C and furin genes are expressed from the mammary gland-specific promoter for whey acidic protein.

ST **transgenic** mammal protein manuf posttranslational modification; mouse **transgenic** protein C furin secretion milk

IT Proteins (specific proteins and subclasses)

RL: BMF (Bioindustrial manufacture); BPN (Biosynthetic preparation); BPR (Biological process); BIOL (Biological study); PREP (Preparation); PROC (Process)

(Gas6 (growth arrest-specific, 6); engineering protein posttranslational modification in **transgenic** non-human mammals)

IT Matrix proteins

RL: BMF (Bioindustrial manufacture); BPN (Biosynthetic preparation); BPR (Biological process); BIOL (Biological study); PREP (Preparation); PROC

(Process)
 (MGP (matrix .gamma.-carboxyglutamic acid-contg. protein); engineering protein posttranslational modification in **transgenic** non-human mammals)

IT Cattle
 Goat
 Guinea pig (*Cavia porcellus*)
 Hamster
 Mammal (Mammalia)
 Mouse
 Post-translational processing
 Rabbit
 Rat
 Sheep
 Swine
 (engineering protein posttranslational modification in **transgenic** non-human mammals)

IT Proteins (general), preparation
 RL: BMF (Bioindustrial manufacture); BPN (Biosynthetic preparation); BIOL (Biological study); PREP (Preparation)
 (engineering protein posttranslational modification in **transgenic** non-human mammals)

IT Albumins, biological studies
 Fibrinogens
 Immunoglobulins
 Osteocalcins
 Protein S (blood coagulation factor)
 RL: BMF (Bioindustrial manufacture); BPN (Biosynthetic preparation); BPR (Biological process); BIOL (Biological study); PREP (Preparation); PROC (Process)
 (engineering protein posttranslational modification in **transgenic** non-human mammals)

IT Mammary gland
 (protein manuf. in; engineering protein posttranslational modification in **transgenic** non-human mammals)

IT Milk
 (protein secretion into; engineering protein posttranslational modification in **transgenic** non-human mammals)

IT Enzymes, biological studies
 RL: BPN (Biosynthetic preparation); BPR (Biological process); BIOL (Biological study); PREP (Preparation); PROC (Process)
 (.gamma.-carboxylating; engineering protein posttranslational modification in **transgenic** non-human mammals)

IT 9000-94-6P, **Antithrombin III** 9001-25-6P,
 Blood-coagulation factor VII 9001-26-7P, Prothrombin 9001-28-9P,
 Blood-coagulation factor IX 9001-29-0P, Blood-coagulation factor X
 11096-26-7P, Erythropoietin 60202-16-6P, Protein C 113189-02-9P,
 Blood-coagulation factor VIII 139639-23-9P, Tissue-type plasminogen activator
 RL: BMF (Bioindustrial manufacture); BPN (Biosynthetic preparation); BPR (Biological process); BIOL (Biological study); PREP (Preparation); PROC (Process)
 (engineering protein posttranslational modification in **transgenic** non-human mammals)

IT 141760-45-4P, **Furin**
 RL: BPN (Biosynthetic preparation); BPR (Biological process); BIOL (Biological study); PREP (Preparation); PROC (Process)
 (engineering protein posttranslational modification in **transgenic** non-human mammals)

IT 246038-41-5 246038-42-6
 RL: PRP (Properties)
 (unclaimed nucleotide sequence; engineering protein posttranslational modification in **transgenic** non-human mammals)

IT 246038-43-7
 RL: PRP (Properties)
 (unclaimed protein sequence; engineering protein posttranslational

modification in **transgenic** non-human mammals)
IT. 245724-26-9 245724-27-0 245724-28-1 245724-29-2

RL: PRP (Properties)

(unclaimed sequence; engineering protein posttranslational
modification

L3 ANSWER 8 OF 24 CAPLUS COPYRIGHT 1999 ACS
 AN 1998:75490 CAPLUS
 DN 128:166377
 TI Commercialization of proteins produced in the mammary gland
 AU Ziomek, C. A.
 CS Genzyme Transgenics Corporation (GTC), Framingham, MA, USA
 SO Theriogenology (1998), 49(1), 139-144
 CODEN: THGNBO; ISSN: 0093-691X
 PB Elsevier Science Inc.
 DT Journal; General Review
 LA English
 CC 16-0 (Fermentation and Bioindustrial Chemistry)
 Section cross-reference(s): 3, 13
 AB A review with 14 refs. In the mid 1980's, a few pioneering companies undertook the risk of developing methodologies for the prodn. of complex human therapeutic proteins in the milk of **transgenic** animals. As we approach the end of the 1990's, the prospect of achieving this aim is becoming a reality as the first of these human therapeutic products, **antithrombin III** and alpha-I-antitrypsin are making their way through human clin. trials. It is projected that licensure by the Regulatory agencies and market launch for these **transgenically** produced therapeutics will occur around the year 2000. Although much has already been achieved, addnl. **transgenic** challenges await the basic embryo researcher and practitioner. The biopharming community recognizes the need for addnl. innovative methodologies (such as cloning, sperm sexing and retroviral mediated gene transfer etc.) to overcome the natural biol. barriers and increase the efficiency of **transgenic** dairy animal prodn. and rapid herd expansion.
 ST review protein prodn mammary gland
 IT Mammary gland
 Transformation (genetic method)
 (commercialization of proteins produced in mammary gland)
 IT **Transgenes**
 RL: BSU (Biological study, unclassified); BIOL (Biological study)
 (commercialization of proteins produced in mammary gland)
 IT Proteins (general), preparation
 RL: BMF (Bioindustrial manufacture); BIOL (Biological study); PREP
 (Preparation)
 (the

L3 ANSWER 12 OF 24 CAPLUS COPYRIGHT 1999 ACS
 AN 1997:489900 CAPLUS
 TI Therapeutic proteins from the milk of **transgenic** livestock:
 Biosynthesis, purification and safety issues.
 AU Velander, William H.
 CS Department Chemical Engineering, Virginia Tech, Blacksburg, VA, 24061,
 USA
 SO Book of Abstracts, 214th ACS National Meeting, Las Vegas, NV, September
 7-11 (1997), MEDI-143 Publisher: American Chemical Society, Washington,
 D.
 C.
 CODEN: 64RNAO
 DT Conference; Meeting Abstract
 LA English
 AB The mammary gland of **transgenic** livestock has been shown to be a
 prodigious bioreactor for making complex, recombinant therapeutic
 proteins. Recombinant human Protein C, Factor IX, Factor VIII,
 Fibrinogen, **Antithrombin III**, and Alpha1-antitrypsin
 have been produced in the milk of pigs, goats and sheep. Two of these
 proteins are currently in human clin. trials. The United States Food and
 Drug Administration has issued a "points to consider document" that
 addresses key regulatory issues facing the manuf. and testing of human
 therapeutics derived from **transgenic** animals. As with other
 biologics, issues concerning product safety and efficacy are centered
 about reproducibility of the therapeutic product and pathogen safety.
 Reproducibility is reflected in several examples of genotypic and
 phenotypic stability which have been shown in **transgenic**
 livestock. The potential for zoonotic disease transmission is species
 specific among livestock. The mammary gland has been shown to be capable
 of making a diversity of post-translational modifications. However, rate
 limitations in glycosylation, gamma-carboxylation of glutamic acid,
 propeptide removal and intra-chain proteolytic processing have been shown
 to occur. Unlike many mammalian cell types, immature forms of the
 recombinant protein appear to be freely secreted by mammary epithelial
 cells. Simplified, nonaffinity, scaleable processes can be used to
 purify
 active recombinant proteins from milk, in spite of the presence of
 immature recombinant and endogenous proteins.

L3 ANSWER 13 OF 24 CAPLUS COPYRIGHT 1999 ACS
 AN 1997:413673 CAPLUS
 DN 127:148357
 TI The past, present, and future of **transgenic** bioreactors
 AU Drohan, William N.
 CS J. Holland Laboratory, American Red Cross, Rockville, MD, 20855, USA
 SO Thromb. Haemostasis (1997), 78(1), 543-547
 CODEN: THHADQ; ISSN: 0340-6245
 PB Schattauer
 DT Journal; General Review
 LA English
 CC 16-0 (Fermentation and Bioindustrial Chemistry)
 Section cross-reference(s): 3
 AB A review is given with 57 refs. Hybrid genes can control the
 tissue-specific synthesis of human proteins in **transgenic**
 animals. It is now possible to produce proteins of biomedical value in
 the body fluids or cells of **transgenic** livestock. The 1st
transgenically produced protein, **antithrombin**
III, is now in clin. trials and others will soon follow.
 ST review **transgenic** animal protein bioreactor
 IT Bioreactors

Transformation (genetic method)

(**transgenic** animals and bioreactors, protein prodn.)

. IT Proteins (general), preparation

RL: BPN (Biosynthetic preparation); BIOL (Biological study); PREP
(Preparation)

(**transgenic** animals and bioreactors, protein prodn.)

L6 ANSWER 9 OF 14 MEDLINE
 AN 86104195 MEDLINE
 DN 86104195
 TI Contribution of **monosaccharide** residues in heparin binding to **antithrombin III**.
 AU Atha D H; Lormeau J C; Petitou M; Rosenberg R D; Choay J
 NC PO1-HL33014 (NHLBI)
 SO BIOCHEMISTRY, (1985 Nov 5) 24 (23) 6723-9.
 Journal code: AOG. ISSN: 0006-2960.
 CY United States
 DT Journal; Article; (JOURNAL ARTICLE)
 LA English
 FS Priority Journals
 EM 198605
 AB The importance of 3-O- and 6-O-sulfated glucosamine residues within the heparin octasaccharide iduronic acid(1)----N-acetylglucosamine 6-O-sulfate(2)----glucuronic acid(3)----N-sulfated glucosamine 3,6-di-O-sulfate(4)----iduronic acid 2-O-sulfate(5)----N-sulfated glucosamine 6-O-sulfate(6)----iduronic acid 2-O-sulfate(7)----anhydromannitol 6-O-sulfate(8) was determined by comparing with synthetic tetra- and penta-saccharides its ability to bind human antithrombin. The octasaccharide had an affinity for antithrombin of 1×10^{-8} M (10.2 kcal/mol) measured by intrinsic fluorescence enhancement at 6 degrees C. The synthetic pentasaccharide, consisting of residues 2-6, had an affinity of 3×10^{-8} M (9.6 kcal/mol). The same pentasaccharide, except lacking the 3-O-sulfate on residue 4, had an affinity of 5×10^{-4} M (4.5 kcal/mol) measured by equilibrium dialysis. The tetrasaccharide, consisting of residues 2-5, bound antithrombin with an affinity of 5×10^{-6} M (6.8 kcal/mol). The tetrasaccharide, consisting of residues 3-6, had an affinity of 5×10^{-5} M (5.5 kcal/mol). Since the loss of either the 6-O-sulfated residue 2 or the 3-O-sulfate of residue 4 results in a 4-5 kcal/mol or a 40-50% loss in binding energy of the pentasaccharide, these two residues must be the major contributors to the binding and must be linked to the biologic activity of the octasaccharide.
 CT Check Tags: Animal; Support, U.S. Gov't, P.H.S.
Antithrombin III: IP, isolation & purification
***Antithrombin III: ME, metabolism**
 Calorimetry
 Carbohydrate Sequence
 Cattle
 Heparin: IP, isolation & purification
 *Heparin: ME, metabolism
 Indicators and Reagents
 Intestinal Mucosa: ME, metabolism
 Oligosaccharides: AN, analysis
 *Oligosaccharides: CS, chemical synthesis
 Spectrometry, Fluorescence
 Swine
 RN 9000-94-6 (Antithrombin III); 9005-49-6 (Heparin)
 CN 0 (Indicators and Reagents); 0 (Oligosaccharides)

L6 ANSWER 11 OF 14 MEDLINE
 AN 85157497 MEDLINE
 DN 85157497
 TI Thrombin-inhibitory activity of whale heparin oligosaccharides.
 AU Ototani N; Kodama C; Kikuchi M; Yosizawa Z
 SO JOURNAL OF BIOCHEMISTRY, (1984 Dec) 96 (6) 1695-703.
 Journal code: HIF. ISSN: 0021-924X.
 CY Japan
 DT Journal; Article; (JOURNAL ARTICLE)
 LA English
 FS Priority Journals
 EM 198507
 AB Whale heparin was partially digested with a purified heparinase and the oligosaccharide fractions with 8-20 **monosaccharide** units were isolated from the digest by gel filtration on Sephadex G-50, followed by affinity chromatography on a column of **antithrombin III** immobilized on Sepharose 4B. A marked difference in the inhibitory activity for thrombin in the presence of **antithrombin III** was observed between the high-affinity fractions for **antithrombin III** of octasaccharide approximately hexadecasaccharide and those of octadecasaccharide approximately eicosasaccharide. The disaccharide compositions of these hexadeca-, octadeca-, and eicosasaccharides were analyzed by high-performance liquid chromatography after digestion with a mixture of purified heparitinases 1 and 2 and heparinase. The analytical data indicated that the proportions of trisulfated disaccharide (IdUA(2S)alpha 1----4GlcNS(6S)) and disulfated disaccharide (UA1----4GlcNS(6S)) increased with the manifestation of high thrombin-inhibitory activity, while that of monosulfated disaccharide (UA1----4GlcNS) decreased. The present observations, together with those so far reported, suggest that the presence of the former structural elements, specifically IdUA(2S)alpha 1----4GlcNS(6S), as well as the **antithrombin III**-binding pentasaccharide at the proper positions in the molecules of whale heparin oligosaccharides is essential for the manifestation of high inhibitory activity for thrombin in the presence of **antithrombin III**. The structural bases for the manifestation of the anticoagulant activity of whale and porcine heparins and their oligosaccharides are also discussed.
 CT Check Tags: Animal; Comparative Study; Support, Non-U.S. Gov't
Antithrombin III: PD, pharmacology
 Carbohydrate Sequence
 *Cetacea: ME, metabolism
 Chromatography, High Pressure Liquid: MT, methods
 *Heparin: PD, pharmacology
 Hydrolysis
 Molecular Weight
 Oligosaccharides: IP, isolation & purification
 *Oligosaccharides: PD, pharmacology
 Polysaccharide-Lyases
 Swine
 *Thrombin: AI, antagonists & inhibitors
 *Whales: ME, metabolism
 RN 9000-94-6 (**Antithrombin III**); 9005-49-6 (Heparin)
 CN EC 3.4.21.5 (Thrombin); EC 4.2.2. (Polysaccharide-Lyases); EC 4.2.2.7 (Heparin Lyase); 0 (Oligosaccharides)

L6 ANSWER 13 OF 14 MEDLINE
 AN 82278007 MEDLINE
 DN 82278007
 TI Effects of heparin oligosaccharides with high affinity for
antithrombin III in experimental venous thrombosis.
 AU Thomas D P; Merton R E; Barrowcliffe T W; Thunberg L; Lindahl U
 SO THROMBOSIS AND HAEMOSTASIS, (1982 Jun 28) 47 (3) 244-8.
 Journal code: VQ7. ISSN: 0340-6245.
 CY GERMANY, WEST: Germany, Federal Republic of
 DT Journal; Article; (JOURNAL ARTICLE)
 LA English
 FS Priority Journals
 EM 198212
 AB The in vitro and in vivo characteristics of two oligosaccharide heparin
 fragments have been compared to those of unfractionated mucosal heparin.
 A
 decasaccharide fragment had essentially no activity by APTT or calcium
 thrombin time assays in vitro, but possessed very high specific activity
 by anti-Factor Xa assays. When injected into rabbits at doses of up to 80
 microgram/kg, this fragment was relatively ineffective in impairing
 stasis
 thrombosis despite producing high blood levels by anti-Xa assays. A 16-18
monosaccharide fragment had even higher specific activity (almost
 2000 iu/mg) by chromogenic substrate anti-Xa assay, with minimal activity
 by APTT. When injected in vivo, this fragment gave low blood levels by
 APTT, very high anti-Xa levels, and was more effective in preventing
 thrombosis than the decasaccharide fragment. However, in comparison with
 unfractionated heparin, the 16-18 **monosaccharide** fragment was
 only partially effective in preventing thrombosis, despite producing much
 higher blood levels by anti-Xa assays. It is concluded that the
 high-affinity binding of a heparin fragment to **antithrombin**
III does not by itself impair venous thrombogenesis, and that the
 anti-Factor Xa activity of heparin is only a partial expression of its
 therapeutic potential.
 CT Check Tags: Animal; Human
 ***Antithrombin III: ME, metabolism**
 Dose-Response Relationship, Drug
 Factor X: AI, antagonists & inhibitors
 Heparin: BL, blood
 *Heparin: TU, therapeutic use
 *Oligosaccharides: TU, therapeutic use
 Partial Thromboplastin Time
 Rabbits
 Thrombin Time
 Thrombophlebitis: BL, blood
 Thrombophlebitis: DI, diagnosis
 *Thrombophlebitis: DT, drug therapy
 RN 9000-94-6 (**Antithrombin III**); 9001-29-0 (Factor X); 9005-49-6
 (Heparin)
 CN EC 3.4

L19 ANSWER 1 OF 1 MEDLINE
 AN 91218798 MEDLINE
 DN 91218798
 TI N-acetyl-D-glucosamine is present in cysts and trophozoites of Giardia lamblia and serves as receptor for wheatgerm agglutinin.
 AU Ortega-Barria E; Ward H D; Evans J E; Pereira M E
 CS Division of Geographic Medicine and Infectious Diseases, New England Medical Center, Tufts University School of Medicine, Boston, MA 02111..
 NC AI 121791 (NIAID)
 SO MOLECULAR AND BIOCHEMICAL PARASITOLOGY, (1990 Dec) 43 (2) 151-65.
 Journal code: NOR. ISSN: 0166-6851.
 CY Netherlands
 DT Journal; Article; (JOURNAL ARTICLE)
 LA English
 FS Priority Journals
 EM 199108
 AB Previously, on the basis of lectin binding and glycosidase digestion assays, we have suggested that N-acetyl-D-glucosamine residues (**GlcNAc**) are major structural components of both trophozoites and in vivo cysts of the intestinal parasite Giardia lamblia. In this report we confirm that **GlcNAc** is present both in trophozoites and in vitro cysts as assessed by lectin binding and glycosidase digestion assays, galactosyltransferase labeling, immunochemical analysis using antibodies specific for **GlcNAc** and its beta 1-4 oligomers, and by gas chromatography/mass spectrometry (GC/MS). The results show that wheatgerm agglutinin (WGA) binds specifically to intact trophozoites and in vitro cysts as well as to SDS-PAGE separated proteins. WGA binding to the separated proteins was markedly reduced after their digestion with N-acetyl-beta-D-glucosaminidase, supporting the conclusion that WGA is reacting with terminal beta-linked **GlcNAc** residues. Labeling of trophozoites and cysts by 3H-exogalactosylation with galactosyltransferase further confirmed the presence of terminal **GlcNAc** in both surface and intracellular glycoproteins. The presence of **GlcNAc** is also supported by microfluorometric analysis using antibodies to (**GlcNAc**)₁, (**GlcNAc**)₂, and (**GlcNAc**)₃, which revealed a sugar-inhibitable binding of the antibody to live trophozoites. Finally, the presence of **GlcNAc** in both cysts and trophozoites was unequivocally confirmed by GC/MS analysis of detergent-extracted membranes and of glycoproteins isolated by affinity chromatography on WGA-agarose. GC/MS analysis also revealed mannose (**Man**), N-acetyl-D-galactosamine (**GalNAc**), fucose (**Fuc**), galactose (**Gal**), glucose (**Glc**) and N-acetylneuraminic acid (**NANA**) to be present in cysts. All these sugars were also present in trophozoites, except for **GalNAc**. The glycoproteins isolated by WGA affinity chromatography were 5- to 40-fold enriched in **GlcNAc**, further supporting the conclusion that WGA reacts with **GlcNAc** in Giardia. In summary, the data presented here provide biological and chemical evidence for **GlcNAc** in both cysts and trophozoites of G. lamblia and are consistent with previously published results from this and other laboratories.
 CT Check Tags: Animal; Support, Non-U.S. Gov't; Support, U.S. Gov't, P.H.S.
 Acetylglucosamine: AN, analysis
 *Acetylglucosamine: ME, metabolism
 Cattle
 Colostrum: EN, enzymology
 Electrophoresis, Polyacrylamide Gel
 Galactosyltransferases: ME, metabolism
 Giardia: AN, analysis

Giardia: GD, growth & development
*Giardia: ME, metabolism
Immunochemistry
Mass Fragmentography
*Wheat Germ Agglutinins: ME, metabolism
RN 7512-17-6 (Acetylglucosamine)
CN EC 2.4.1.- (Galactosyltransferases)

L46 ANSWER 1 OF 4 BIOSIS COPYRIGHT 1999 BIOSIS
 AN 1999:69789 BIOSIS
 DN PREV199900069789
 TI **Transgenically** produced **antithrombin III**.
 AU Ditullio, P.; Meade, H.; Cole, E. S.
 CS Framingham, Mass. USA
 ASSIGNEE: GENZYME TRANSGENIC CORPORATION
 PI US 5843705 Dec. 1, 1998
 SO Official Gazette of the United States Patent and Trademark Office
 Patents,
 (Dec. 1, 1998) Vol. 1217, No. 1, pp. 479.
 ISSN: 0098-1133.
 DT Patent
 LA English
 NCL 435069001
 CC *00100
 *04600
 *15100
 *16200
 *21100
 *21300
 *51200
 *51400
 *80100
 IT Major Concepts
 Animal Husbandry (Agriculture); Biochemistry and Molecular Biophysics;
 Blood and Lymphatics (Transport and Circulation); General Life
 Studies;
 Genetics; Miscellaneous Substances; Pharmacology; Reproductive System
 (Reproduction)
 IT Miscellaneous Descriptors
 BIOTECHNOLOGY; ENCODING; GOAT **MILK**; HUMAN
ANTITHROMBIN III; MAMMARY TISSUE; PHARMACEUTICALS;
TRANSGENE EXPRESSION; **TRANSGENIC** GOAT
 ORGN Super Taxa
 Mammalia - Unspecified: Mammalia, Vertebrata, Chordata, Animalia
 ORGN Organism Name
 mammal (Mammalia - Unspecified)
 ORGN Organism Superterms
 animals; chordates; mammals; nonhuman vertebrates; nonhuman mammals;
 vertebrates

 L46 ANSWER 2 OF 4 BIOSIS COPYRIGHT 1999 BIOSIS
 AN 1997:428755 BIOSIS
 DN PREV199799727958
 TI Therapeutic proteins from the **milk** of **transgenic**
 livestock: Biosynthesis, purification and safety issues.
 AU Velander, William H.
 CS Dep. Chem. Engineering, Virginia Tech, Blacksburg, VA 24061 USA
 SO Abstracts of Papers American Chemical Society, (1997) Vol. 214, No. 1-2,
 pp. MEDI 143.
 Meeting Info.: 214th American Chemical Society National Meeting Las
 Vegas,
 Nevada, USA September 7-11, 1997
 ISSN: 0065-7727.
 DT Conference; Abstract
 LA English
 CC General Biology - Symposia, Transactions and Proceedings of Conferences,
 Congresses, Review Annuals 00520
 Cytology and Cytochemistry - Animal *02506

Genetics and Cytogenetics - Animal *03506
 Comparative Biochemistry, General *10010
 Biochemical Methods - Nucleic Acids, Purines and Pyrimidines *10052
 Biochemical Methods - Proteins, Peptides and Amino Acids *10054
 Biochemical Studies - Proteins, Peptides and Amino Acids *10064
 Biophysics - Molecular Properties and Macromolecules *10506
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 Blood, Blood-Forming Organs and Body Fluids - Other Body Fluids *15010
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 Endocrine System - General *17002
 Pharmacology - General *22002
 Animal Production - General; Methods *26502
 BC Mammalia - Unspecified 85700
 Bovidae 85715
 Suidae *85740
 IT Major Concepts
 Animal Husbandry (Agriculture); Biochemistry and Molecular Biophysics;
 Blood and Lymphatics (Transport and Circulation); Cell Biology;
 Endocrine System (Chemical Coordination and Homeostasis); Genetics;
 Metabolism; Methods and Techniques; Pathology; Pharmacology;
 Physiology; Reproductive System (Reproduction)
 IT Chemicals & Biochemicals
 PROTEIN C; FACTOR VII; **ANTITHROMBIN III**;
 ALPHA-1-ANTITRYPSIN
 IT Miscellaneous Descriptors
 ALPHA-1-ANTITRYPSIN; ANIMAL HUSBANDRY; **ANTITHROMBIN**
 III; BIOSYNTHESIS; BIOTECHNOLOGY; FACTOR IX; FACTOR VII;
 FIBRINOGEN; GENOTYPE; MAMMARY GLAND; **MILK**; MOLECULAR
 GENETICS; PHARMACOLOGY; PHENOTYPE; PROTEIN C; PROTEINS; PROTEOLYTIC
 PROCESSING; PURIFICATION; RECOMBINANT PROTEINS; REPRODUCTIVE SYSTEM;
 SAFETY; THERAPEUTIC PROTEINS; **TRANSGENIC**
 ORGN Super Taxa
 Bovidae: Artiodactyla, Mammalia, Vertebrata, Chordata, Animalia;
 Mammalia - Unspecified: Mammalia, Vertebrata, Chordata, Animalia;
 Suidae: Artiodactyla, Mammalia, Vertebrata, Chordata, Animalia
 ORGN Organism Name
 goat (Bovidae); livestock (Mammalia - Unspecified); pig (Suidae);
 sheep
 (Bovidae); Mammalia (Mammalia - Unspecified)
 ORGN Organism Superterms
 animals; artiodactyls; chordates; mammals; nonhuman mammals; nonhuman
 vertebrates; vertebrates
 RN 60202-16-6 (PROTEIN C)
 9001-25-6 (FACTOR VII)
 9000-94-6 (**ANTITHROMBIN III**)
 9041-92-3 (ALPHA-1-ANTITRYPSIN)

 L46 ANSWER 3 OF 4 BIOSIS COPYRIGHT 1999 BIOSIS
 AN 1994:250575 BIOSIS
 DN PREV199497263575
 TI Tissue specific and species differences in the glycosylation pattern of
 antithrombin III.
 AU Edmunds, Tim; Higgins, Elizabeth; Bernasconi, Rick; Garone, Louise; Cole,
 Edward S.
 CS GENZYME Corp., 1 Mountain Road, Framingham, MA 01701-9322 USA
 SO Journal of Cellular Biochemistry Supplement, (1994) Vol. 0, No. 18D, pp.
 265.
 Meeting Info.: Keystone Symposium on Complex Carbohydrates in Biology and
 Medicine Frisco, Colorado, USA March 19-26, 1994
 ISSN: 0733-1959.
 DT Conference
 LA English
 CC General Biology - Symposia, Transactions and Proceedings of Conferences,

Congresses, Review Annuals 00520
 Genetics and Cytogenetics - Animal *03506
 Genetics and Cytogenetics - Human *03508
 Comparative Biochemistry, General *10010
 Biochemical Methods - Carbohydrates 10058
 Biochemical Studies - Proteins, Peptides and Amino Acids *10064
 Biochemical Studies - Carbohydrates *10068
 Biophysics - General Biophysical Techniques 10504
 Biophysics - Molecular Properties and Macromolecules *10506
 Metabolism - Carbohydrates *13004
 Metabolism - Proteins, Peptides and Amino Acids *13012
 Blood, Blood-Forming Organs and Body Fluids - Blood and Lymph Studies
 *15002
 Reproductive System - Physiology and Biochemistry *16504
 Immunology and Immunochemistry - Immunopathology, Tissue Immunology
 *34508
 BC Bovidae 85715
 Hominidae *86215
 IT Major Concepts
 Biochemistry and Molecular Biophysics; Blood and Lymphatics (Transport
 and Circulation); Clinical Immunology (Human Medicine, Medical
 Sciences); Genetics; Metabolism; Reproductive System (Reproduction)
 IT Chemicals & Biochemicals
ANTITHROMBIN III
 IT Miscellaneous Descriptors
 FLUOROPHORE ASSISTED CARBOHYDRATE ANALYSIS; MEETING ABSTRACT; MEETING
 POSTER; **MILK**; **TRANSGENIC** GOATS
 ORGN Super Taxa
 Bovidae: Artiodactyla, Mammalia, Vertebrata, Chordata, Animalia;
 Hominidae: Primates, Mammalia, Vertebrata, Chordata, Animalia
 ORGN Organism Name
 human (Hominidae); Bovidae (Bovidae)
 ORGN Organism Superterms
 animals; artiodactyls; chordates; humans; mammals; nonhuman mammals;
 nonhuman vertebrates; primates; vertebrates
 RN 9000-94-6 (**ANTITHROMBIN III**)
 L46 ANSWER 4 OF 4 BIOSIS COPYRIGHT 1999 BIOSIS
 AN 1994:250573 BIOSIS
 DN PREV199497263573
 TI Glycosylation patterns of human proteins expressed in **transgenic**
 goat **milk**.
 AU Cole, Edward S.; Higgins, Elizabeth; Bernasconi, Rick; Garone, Louise;
 Edmunds, Tim
 CS GENZYME Corp., 1 Mountain Road, Framingham, MA 01701-9322 USA
 SO Journal of Cellular Biochemistry Supplement, (1994) Vol. 0, No. 18D, pp.
 265.
 Meeting Info.: Keystone Symposium on Complex Carbohydrates in Biology and
 Medicine Frisco, Colorado, USA March 19-26, 1994
 ISSN: 0733-1959.
 DT Conference
 LA English
 CC General Biology - Symposia, Transactions and Proceedings of Conferences,
 Congresses, Review Annuals 00520
 Genetics and Cytogenetics - Animal *03506
 Genetics and Cytogenetics - Human *03508
 Comparative Biochemistry, General *10010
 Biochemical Studies - Nucleic Acids, Purines and Pyrimidines 10062
 Biochemical Studies - Proteins, Peptides and Amino Acids *10064
 Biochemical Studies - Carbohydrates *10068
 Replication, Transcription, Translation 10300
 Biophysics - General Biophysical Techniques 10504
 Biophysics - Molecular Properties and Macromolecules *10506
 Metabolism - Carbohydrates *13004
 Metabolism - Proteins, Peptides and Amino Acids *13012
 Blood, Blood-Forming Organs and Body Fluids - Other Body Fluids *15010

Reproductive System - Physiology and Biochemistry *16504

BC Bovidae 85715
 Hominidae *86215

IT Major Concepts
 Biochemistry and Molecular Biophysics; Genetics; Metabolism;
 Physiology; Reproductive System (Reproduction)

IT Chemicals & Biochemicals
 N-ACETYLNEURAMINIC ACID; **ANTITHROMBIN-III**

IT Miscellaneous Descriptors
ANTITHROMBIN-III, TISSUE PLASMINOGEN ACTIVATOR;
 COMPLEX CARBOHYDRATES; ELECTROSPRAY MASS SPECTROMETRY; FLUOROPHORE
 ASSISTED CARBOHYDRATE ELECTROPHORESIS; MEETING ABSTRACT; MEETING
 POSTER; N-ACETYLNEURAMINIC ACID; N-GLYCOLYLNEURAMINIC ACID

ORGN Super Taxa
 Bovidae: Artiodactyla, Mammalia, Vertebrata, Chordata, Animalia;
 Hominidae: Primates, Mammalia, Vertebrata, Chordata, Animalia

ORGN Organism Name
 Bovidae (Bovidae); Hominidae (Hominidae)

ORGN Organism Superterms
 animals; artiodactyls; chordates; humans; mammals; nonhuman mammals;
 nonhuman vertebrates; primates; vertebrates

RN 131-48-6 (N-ACETYLNEURAMINIC ACID)
 9000-94-6 (**ANTITHROMBIN-III**)

L8 ANSWER 1 OF 1 CAPLUS COPYRIGHT 1999 ACS

AN 1997:343782 CAPLUS

DN 127:61135

TI Adrenomedullin as an autocrine/paracrine apoptosis survival factor for
rat endothelial cells

AU Kato, Hiroki; Shichiri, Masayoshi; Marumo, Fumiaki; Hirata, Yukio

CS Endocrine-Hypertension Division, Second Department of Internal Medicine,
Tokyo Medical and Dental University, Tokyo, 113, Japan

SO Endocrinology (1997), 138(6), 2615-2620

CODEN: ENDOAO; ISSN: 0013-7227

PB Endocrine Society

DT Journal

LA English

CC 2-10 (Mammalian Hormones)

AB Adrenomedullin is a potent vasorelaxant/hypotensive peptide recently
isolated from human pheochromocytoma. We demonstrate here a novel role
of

this peptide as an apoptosis survival factor for rat endothelial cells.
When rendered quiescent by serum deprivation, a fraction of endothelial
cell cultures showed morphol. and biochem. features characteristic of
apoptosis. Adrenomedullin significantly suppressed apoptosis without
inducing cell proliferation. Rat endothelial cells that contained
high affinity binding sites for adrenomedullin
expressed adrenomedullin **gene** and released the peptide into
culture media. Addn. of preimmune rabbit serum prevented apoptosis,
whereas rabbit antiadrenomedullin antiserum partially, but significantly,
abrogated the protective effect of the preimmune serum, suggesting its
autocrine/paracrine role. Although adrenomedullin induced intracellular
cAMP formation, other cAMP-elevating agonists, such as prostaglandin I₂
and forskolin, did not affect apoptosis. Furthermore, adenosine
3',5'-cyclic monophosphothioate Rp-isomer, a cAMP antagonist, did not
block the cell survival effect of adrenomedullin. Adrenomedullin neither
increased intracellular Ca²⁺ concns. nor **inositol-1,**
4,5-trisphosphate levels in rat endothelial
cells. These results demonstrate that adrenomedullin suppresses serum
deprivation-induced apoptosis of rat endothelial cells via
cAMP-independent mechanism.

ST adrenomedullin apoptosis **vascular** endothelium cAMP

IT Apoptosis

Calcium transport (biological)

Cell proliferation

Second messenger system

Serum (blood)

Vascular endothelium

(adrenomedullin suppresses serum deprivation-induced apoptosis of
endothelial cells via cAMP-independent mechanism)

IT Genes (animal)

RL: BPR (Biological process); BIOL (Biological study); PROC (Process)

(adrenomedullin suppresses serum deprivation-induced apoptosis of
endothelial cells via cAMP-independent mechanism)

IT Hormone receptors

RL: BPR (Biological process); BIOL (Biological study); PROC (Process)

(adrenomedullin; adrenomedullin suppresses serum deprivation-induced
apoptosis of endothelial cells via cAMP-independent mechanism)

IT 154835-90-2, Adrenomedullin

RL: BAC (Biological activity or effector, except adverse); BIOL
(Biological study)

(adrenomedullin suppresses serum deprivation-induced apoptosis of
endothelial cells via cAMP-independent mechanism)

IT 60-92-4, CAMP
RL: BAC (Biological activity or effector, except adverse); BPR
(Biological
process); BIOL (Biological study); PROC (Process)
(adrenomedullin suppresses serum deprivation-induced apoptosis of
endothelial cells via cAMP-independent mechanism)

IT 7440-70-2, Calcium, biological studies 88269-39-0, **Inositol-
1,4,5-trisphosphate**
RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
(adrenomedullin suppresses serum deprivation-induced apoptosis of
endothelial cells via cAMP-independent mechanism)

6 ANSWER 1 OF 1 CAPLUS COPYRIGHT 1999 ACS

AN 1999:343182 CAPLUS

DN 131:140289

TI Molecular cloning of the mouse follicle-stimulating hormone receptor complementary deoxyribonucleic acid: functional expression of alternatively spliced variants and receptor inactivation by a C566T transition in Exon 7 of the coding sequence

AU Tena-Sempere, Manuel; Manna, Pulak R.; Huhtaniemi, Ilpo

CS Department of Physiology, University of Turku, Turku, 20520, Finland

SO Biol. Reprod. (1999), 60(6), 1515-1527

CODEN: BIREBV; ISSN: 0006-3363

PB Society for the Study of Reproduction

DT Journal

LA English

CC 3-3 (Biochemical Genetics)

Section cross-reference(s): 2, 14

AB The gonadotropin receptors, i.e., those of LH and FSH (FSHR), are pivotal elements in the regulation of gonadal function. Recently, extensive efforts have been made to elucidate the structure-function relation of these receptors as well as the modulatory mechanism(s) of their function. In the present study, the authors report (1) characterization of the

mouse

(m) FSHR cDNA coding sequence and (2) the functional consequences of coexpression of several splice variants of the mFSHR. In addn., the authors evaluate (3) the impact on mFSHR function of a C566T transition

in

exon 7 of the coding sequence, a substitution analogous to the inactivating mutation in the human FSHR **gene** responsible for a hereditary form of hypergonadotropic ovarian failure. Mol. cloning of

the

mFSHR cDNA was carried out by reverse transcription-polymerase chain reaction (RT-PCR) using 129/Sv mouse testicular RNA and primers complementary to the rat or the partially characterized mouse FSHR sequence. Overlapping partial fragments of receptor cDNA were amplified, sequenced, and engineered to produce the entire cDNA coding sequence, subcloned into the pSG5 expression vector. Using a similar approach, 4 different receptor splice variants, selectively lacking exons 2, 2 and 5, 5 and 6, and 2, 5, and 6 of the coding region, were cloned. Finally, PCR-based site-directed mutagenesis was used to generate the C566T mutant of mFSHR. Sequence anal. showed an open reading frame of 2076 base pairs for the mFSHR cDNA, predicting a putative 17-amino acid signal peptide

and

a 675-amino acid mature receptor protein, and overall sequence homol. of 94% with rat, 87% with human, and 85-84% with bovine, and ovine FSHRs. Functional expression in human embryonic kidney (HEK 293) and mouse granulosa (KK-1) cells demonstrated for the cloned receptor **high -affinity binding** to recombinant human (rh) FSH and ability to elicit cAMP, **inositol** trisphosphate (IP3), and progesterone responses. In contrast, transient transfection studies showed that despite successful transcription, the exon-lacking FSHR variants were unable to bind rhFSH either in intact or in solubilized HEK 293 cells, or to elicit cAMP or progesterone responses in KK-1 cells. Furthermore, cotransfections of the splice variants in the context of an ovarian cell line stably expressing the full-length mFSHR failed to demonstrate modulatory effects on the holoreceptor function. Finally, transient expression of the C566T mFSHR mutant in HEK 293 cells revealed that, in accordance with observations on human FSHR, this substitution profoundly impaired the ligand binding and cAMP and IP3 responses to

rhFSH

stimulation. In conclusion, the present data indicate that, despite

extensive splicing of the mFSHR message, a potential role of the exon-lacking receptor transcripts in modulating FSH actions is unlikely. In addn., the authors provide evidence for mFSHR inactivation by a C566T transition in exon 7 of the coding sequence, thus paving the way for further development of animal models of hypergonadotropic ovarian failure.

ST mouse FSH receptor cDNA sequence expression hypergonadotropic ovarian failure; **gene** FSH receptor mutation hypergonadotropic ovarian failure; splicing variant FSH receptor expression

IT Genes (animal)
 RL: BOC (Biological occurrence); PRP (Properties); BIOL (Biological study); OCCU (Occurrence)
 (Fshr; sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT Ovarian **diseases**
 (failure, hypergonadotropic; sequence of mouse FSH receptor cDNA, expression of spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT Protein sequences
 (homol.; protein sequence homol. of mouse FSH receptor with other mammalian FSH receptors)

IT Splicing (RNA)
 (messenger; sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by C566T transition in exon 7 of coding sequence)

IT **Disease** models
Gene expression
 Mouse (Mus musculus)
 Protein sequences
 Transcription (genetic)
 Transition mutation
 cDNA sequences
 (sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT Exon (genetic element)
 RL: ADV (Adverse effect, including toxicity); BOC (Biological occurrence);
 PRP (Properties); BIOL (Biological study); OCCU (Occurrence)
 (sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT mRNA
 RL: BOC (Biological occurrence); BPR (Biological process); MFM (Metabolic formation); PRP (Properties); BIOL (Biological study); FORM (Formation, nonpreparative); OCCU (Occurrence); PROC (Process)
 (sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT FSH receptors
 RL: BOC (Biological occurrence); BPR (Biological process); PRP (Properties); BIOL (Biological study); OCCU (Occurrence); PROC (Process)
 (sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT Pre-mRNA
 RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
 (splicing; sequence of mouse FSH receptor cDNA, functional expression

of alternatively spliced variants in human and mouse cells and
receptor
inactivation by C566T transition in exon 7 of coding sequence)

IT 57-83-0, Progesterone, biological studies 60-92-4, CAMP 85166-31-0,
D-myo-Inositol 1,4,5-
trisphosphate
RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
(FSH-induced response; sequence of mouse FSH receptor cDNA, functional
expression of alternatively spliced variants in human and mouse cells
and receptor inactivation by C566T transition in exon 7 of coding
sequence)

IT 234764-60-4
RL: PRP (Properties)
(amino acid sequence; sequence of mouse FSH receptor cDNA, functional
expression of alternatively spliced variants in human and mouse cells
and receptor inactivation by C566T transition in exon 7 of coding
sequence)

IT 218721-47-2, GenBank AF095642
RL: PRP (Properties)
(nucleotide sequence; sequence of mouse FSH receptor cDNA, functional
expression of alternatively spliced variants in human and mouse cells
and receptor inactivation by C566T transition in exon 7 of coding
sequence)

IT 9002-68-0, FSH
RL: BAC (Biological activity or effector, except adverse); BPR
(Biological
process); BIOL (Biological study); PROC (Process)
(sequence of mouse FSH receptor cDNA, functional expression of
alternatively spliced variants in human and mouse cells and receptor
inactivation by hypergonadotropic ovarian failure-assocd. C566T
transition in exon 7 of coding sequence)

L5 ANSWER 1 OF 8 CAPLUS COPYRIGHT 1999 ACS
 AN 1999:537292 CAPLUS
 DN 131:346963
 TI Free-radical-generated F2-isoprostane stimulates cell proliferation and endothelin-1 expression on endothelial cells
 AU Yura, Takafumi; Fukunaga, Megumu; Khan, Rizwan; Nassar, George N.; Badr, Kamal F.; Montero, Angel
 CS Renal Division, Department of Medicine, Emory University and Veterans Affairs Medical Center, Atlanta, GA, USA
 SO Kidney Int. (1999), 56(2), 471-478
 CODEN: KDYIA5; ISSN: 0085-2538
 PB Blackwell Science, Inc.
 DT Journal
 LA English
 CC 2-9 (Mammalian Hormones)
 AB Free-radical-generated F2-isoprostane stimulates DNA synthesis and endothelin-1 (ET-1) expression on endothelial cells. 8-Iso-prostaglandin F2.alpha. (8-iso-PGF2.alpha.) is a member of the recently discovered family of prostanoids, the F2-isoprostanes, produced in vivo by cyclooxygenase-independent, free-radical-catalyzed lipid peroxidn. The goal of the authors' study is to establish the effect of isoprostane on ET-1 prodn. by endothelial cells, as well to det. the receptors responsible for these effects. The proliferative effect of isoprostanes was measured as an increase of viable cell no. and [3H]-thymidine uptake. ET-1 **gene** expression and protein synthesis were detd. by Northern blot and RIA, resp. The authors also detd. **inositol 1,4,5-trisphosphate** synthesis. Thromboxane A2 (TXA2) receptor antagonist SQ29,548 was used to establish the role of TXA2 receptor in isoprostane effect, as well as to det. the type of receptors involved in these effects. The authors' results show that physiol. concns. of 8-iso-PGF2.alpha. stimulated cell proliferation, DNA synthesis, and ET-1 mRNA and protein expression in bovine aortic endothelial cells (BAECs). The proliferative effect was partially abolished by treatment with anti-endothelin antibody. 8-Iso-PGF2.alpha. also increased **inositol 1,4,5-trisphosphate** formation in these cells. These effects were partially inhibited by SQ29,548. In competitive binding assays, two binding sites were recognized on BAECs with dissocn. consts. (Kd) and binding site densities at equil. similar to those previously described in smooth muscle cells and likely represent [3H]-8-iso-PGF2.alpha. binding to its own receptor (**high-affinity binding site**) and cross-recognition of the TXA2 receptor (**low-affinity binding site**). These studies expand the potential scope of the pathophysiol. significance of F2-isoprostanes, released during oxidant injury, to include alteration of endothelial cell biol.
 ST PGF 2alpha ET1 expression proliferation endothelium signaling free radical
 IT Artery
 (aorta, endothelium, bovine; free radical generated F2-isoprostane stimulates cell proliferation and endothelin-1 expression on endothelial cells)
 IT Cell proliferation
 Radical ions
 Second messenger system
 Tobacco smoke
 Transcriptional regulation
 (free radical generated F2-isoprostane stimulates cell proliferation and endothelin-1 expression on endothelial cells)
 IT Thromboxane receptors

RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
 (free radical generated F2-isoprostane stimulates cell proliferation
 and endothelin-1 expression on endothelial cells)

IT Phosphoinositides
 RL: BPR (Biological process); BSU (Biological study, unclassified); BIOL
 (Biological study); PROC (Process)
 (free radical generated F2-isoprostane stimulates cell proliferation
 and endothelin-1 expression on endothelial cells)

IT 27415-26-5, 8-Iso-prostaglandin F2.alpha.
 RL: BAC (Biological activity or effector, except adverse); BIOL
 (Biological study)
 (free radical generated F2-isoprostane stimulates cell proliferation
 and endothelin-1 expression on endothelial cells)

IT 123626-67-5, Endothelin-1
 RL: BPR (Biological process); MFM (Metabolic formation); BIOL (Biological
 study); FORM (Formation, nonpreparative); PROC (Process)
 (free radical generated F2-isoprostane stimulates cell proliferation
 and endothelin-1 expression on endothelial cells)

IT 88269-39-0, **Inositol-1,4,5-**
trisphosphate
 RL: MFM (Metabolic formation); BIOL (Biological study); FORM (Formation,
 nonpreparative)
 (free radical generated F2-isoprostane stimulates cell proliferation
 and endothelin-1 expression on endothelial cells)

L5 ANSWER 2 OF 8 CAPLUS COPYRIGHT 1999 ACS

AN 1999:343182 CAPLUS

DN 131:140289

TI Molecular cloning of the mouse follicle-stimulating hormone receptor
 complementary deoxyribonucleic acid: functional expression of
 alternatively spliced variants and receptor inactivation by a C566T
 transition in Exon 7 of the coding sequence

AU Tena-Sempere, Manuel; Manna, Pulak R.; Huhtaniemi, Ilpo

CS Department of Physiology, University of Turku, Turku, 20520, Finland

SO Biol. Reprod. (1999), 60(6), 1515-1527

CODEN: BIREBV; ISSN: 0006-3363

PB Society for the Study of Reproduction

DT Journal

LA English

CC 3-3 (Biochemical Genetics)

Section cross-reference(s): 2, 14

AB The gonadotropin receptors, i.e., those of LH and FSH (FSHR), are pivotal
 elements in the regulation of gonadal function. Recently, extensive
 efforts have been made to elucidate the structure-function relation of
 these receptors as well as the modulatory mechanism(s) of their function.
 In the present study, the authors report (1) characterization of the

mouse

(m) FSHR cDNA coding sequence and (2) the functional consequences of
 coexpression of several splice variants of the mFSHR. In addn., the
 authors evaluate (3) the impact on mFSHR function of a C566T transition

in

exon 7 of the coding sequence, a substitution analogous to the
 inactivating mutation in the human FSHR **gene** responsible for a
 hereditary form of hypergonadotropic ovarian failure. Mol. cloning of

the

mFSHR cDNA was carried out by reverse transcription-polymerase chain
 reaction (RT-PCR) using 129/Sv mouse testicular RNA and primers
 complementary to the rat or the partially characterized mouse FSHR
 sequence. Overlapping partial fragments of receptor cDNA were amplified,
 sequenced, and engineered to produce the entire cDNA coding sequence,
 subcloned into the pSG5 expression vector. Using a similar approach, 4
 different receptor splice variants, selectively lacking exons 2, 2 and 5,
 5 and 6, and 2, 5, and 6 of the coding region, were cloned. Finally,
 PCR-based site-directed mutagenesis was used to generate the C566T mutant
 of mFSHR. Sequence anal. showed an open reading frame of 2076 base pairs
 for the mFSHR cDNA, predicting a putative 17-amino acid signal peptide

and

a 675-amino acid mature receptor protein, and overall sequence homol. of 94% with rat, 87% with human, and 85-84% with bovine, and ovine FSHRs. Functional expression in human embryonic kidney (HEK 293) and mouse granulosa (KK-1) cells demonstrated for the cloned receptor **high-affinity binding** to recombinant human (rh) FSH and ability to elicit cAMP, **inositol** trisphosphate (IP3), and progesterone responses. In contrast, transient transfection studies showed that despite successful transcription, the exon-lacking FSHR variants were unable to bind rhFSH either in intact or in solubilized HEK 293 cells, or to elicit cAMP or progesterone responses in KK-1 cells. Furthermore, cotransfections of the splice variants in the context of an ovarian cell line stably expressing the full-length mFSHR failed to demonstrate modulatory effects on the holoreceptor function. Finally, transient expression of the C566T mFSHR mutant in HEK 293 cells revealed that, in accordance with observations on human FSHR, this substitution profoundly impaired the ligand binding and cAMP and IP3 responses to rhFSH stimulation. In conclusion, the present data indicate that, despite extensive splicing of the mFSHR message, a potential role of the exon-lacking receptor transcripts in modulating FSH actions is unlikely. In addn., the authors provide evidence for mFSHR inactivation by a C566T transition in exon 7 of the coding sequence, thus paving the way for further development of animal models of hypergonadotropic ovarian failure.

ST mouse FSH receptor cDNA sequence expression hypergonadotropic ovarian failure; **gene** FSH receptor mutation hypergonadotropic ovarian failure; splicing variant FSH receptor expression

IT Genes (animal)
 RL: BOC (Biological occurrence); PRP (Properties); BIOL (Biological study); OCCU (Occurrence)
 (Fshr; sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT Ovarian diseases
 (failure, hypergonadotropic; sequence of mouse FSH receptor cDNA, expression of spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT Protein sequences
 (homol.; protein sequence homol. of mouse FSH receptor with other mammalian FSH receptors)

IT Splicing (RNA)
 (messenger; sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by C566T transition in exon 7 of coding sequence)

IT Disease models
Gene expression
 Mouse (Mus musculus)
 Protein sequences
 Transcription (genetic)
 Transition mutation
 cDNA sequences
 (sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT Exon (genetic element)
 RL: ADV (Adverse effect, including toxicity); BOC (Biological occurrence);
 PRP (Properties); BIOL (Biological study); OCCU (Occurrence)
 (sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT mRNA
 RL: BOC (Biological occurrence); BPR (Biological process); MFM (Metabolic formation); PRP (Properties); BIOL (Biological study); FORM (Formation, nonpreparative); OCCU (Occurrence); PROC (Process)
 (sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT FSH receptors
 RL: BOC (Biological occurrence); BPR (Biological process); PRP (Properties); BIOL (Biological study); OCCU (Occurrence); PROC (Process)
 (sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

IT Pre-mRNA
 RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
 (splicing; sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by C566T transition in exon 7 of coding sequence)

IT 57-83-0, Progesterone, biological studies 60-92-4, CAMP 85166-31-0, D-myo-Inositol 1,4,5-trisphosphate
 RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
 (FSH-induced response; sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by C566T transition in exon 7 of coding sequence)

IT 234764-60-4
 RL: PRP (Properties)
 (amino acid sequence; sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by C566T transition in exon 7 of coding sequence)

IT 218721-47-2, GenBank AF095642
 RL: PRP (Properties)
 (nucleotide sequence; sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by C566T transition in exon 7 of coding sequence)

IT 9002-68-0, FSH
 RL: BAC (Biological activity or effector, except adverse); BPR (Biological process); BIOL (Biological study); PROC (Process)
 (sequence of mouse FSH receptor cDNA, functional expression of alternatively spliced variants in human and mouse cells and receptor inactivation by hypergonadotropic ovarian failure-assocd. C566T transition in exon 7 of coding sequence)

L5 ANSWER 3 OF 8 CAPLUS COPYRIGHT 1999 ACS
 AN 1997:343782 CAPLUS
 DN 127:61135
 TI Adrenomedullin as an autocrine/paracrine apoptosis survival factor for rat endothelial cells
 AU Kato, Hiroki; Shichiri, Masayoshi; Marumo, Fumiaki; Hirata, Yukio
 CS Endocrine-Hypertension Division, Second Department of Internal Medicine, Tokyo Medical and Dental University, Tokyo, 113, Japan
 SO Endocrinology (1997), 138(6), 2615-2620
 CODEN: ENDOAO; ISSN: 0013-7227
 PB Endocrine Society
 DT Journal
 LA English
 CC 2-10 (Mammalian Hormones)
 AB Adrenomedullin is a potent vasorelaxant/hypotensive peptide recently

isolated from human pheochromocytoma. We demonstrate here a novel role of this peptide as an apoptosis survival factor for rat endothelial cells. When rendered quiescent by serum deprivation, a fraction of endothelial cell cultures showed morphol. and biochem. features characteristic of apoptosis. Adrenomedullin significantly suppressed apoptosis without inducing cell proliferation. Rat endothelial cells that contained **high affinity binding** sites for adrenomedullin expressed adrenomedullin **gene** and released the peptide into culture media. Addn. of preimmune rabbit serum prevented apoptosis, whereas rabbit antiadrenomedullin antiserum partially, but significantly, abrogated the protective effect of the preimmune serum, suggesting its autocrine/paracrine role. Although adrenomedullin induced intracellular cAMP formation, other cAMP-elevating agonists, such as prostaglandin I2 and forskolin, did not affect apoptosis. Furthermore, adenosine 3',5'-cyclic monophosphothioate Rp-isomer, a cAMP antagonist, did not block the cell survival effect of adrenomedullin. Adrenomedullin neither increased intracellular Ca²⁺ concns. nor **inositol-1, 4,5-trisphosphate** levels in rat endothelial cells. These results demonstrate that adrenomedullin suppresses serum deprivation-induced apoptosis of rat endothelial cells via cAMP-independent mechanism.

ST adrenomedullin apoptosis vascular endothelium cAMP
IT Apoptosis
Calcium transport (biological)
Cell proliferation
Second messenger system
Serum (blood)
Vascular endothelium
(adrenomedullin suppresses serum deprivation-induced apoptosis of endothelial cells via cAMP-independent mechanism)

IT Genes (animal)
RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
(adrenomedullin suppresses serum deprivation-induced apoptosis of endothelial cells via cAMP-independent mechanism)

IT Hormone receptors
RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
(adrenomedullin; adrenomedullin suppresses serum deprivation-induced apoptosis of endothelial cells via cAMP-independent mechanism)

IT 154835-90-2, Adrenomedullin
RL: BAC (Biological activity or effector, except adverse); BIOL (Biological study)
(adrenomedullin suppresses serum deprivation-induced apoptosis of endothelial cells via cAMP-independent mechanism)

IT 60-92-4, CAMP
RL: BAC (Biological activity or effector, except adverse); BPR (Biological process); BIOL (Biological study); PROC (Process)
(adrenomedullin suppresses serum deprivation-induced apoptosis of endothelial cells via cAMP-independent mechanism)

IT 7440-70-2, Calcium, biological studies 88269-39-0, **Inositol-1,4,5-trisphosphate**
RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
(adrenomedullin suppresses serum deprivation-induced apoptosis of endothelial cells via cAMP-independent mechanism)

L5 ANSWER 4 OF 8 CAPLUS COPYRIGHT 1999 ACS
AN 1995:767118 CAPLUS
DN 124:22062
TI Selection and characterization of mammalian cell lines with stable over-expression of human pituitary receptors for gonadoliberin
AU Beckers, Thomas; Marheineke, Kathrin; Reilaender, Helmut; Hilgard, Peter
CS ASTA Medica AG, Frankfurt/Main, Germany
SO Eur. J. Biochem. (1995), 231(3), 535-43
CODEN: EJBCAI; ISSN: 0014-2956
DT Journal

LA English
CC 2-5 (Mammalian Hormones)
AB The cDNA encoding the receptor for gonadoliberein (GnRH or LH-RH) was isolated from a human pituitary cDNA library and heterologously expressed in the murine fibroblast cell line LTK-. By using a dicistronic expression strategy utilizing the internal ribosomal-entry-site sequence of poliovirus, single cell clones with stable and high expression of human gonadoliberein receptors were selected. The gonadoliberein antagonist Cetorelix showed **high-affinity binding** to the heterologously expressed human gonadoliberein receptor with a Kd of 0.1 nM in radioligand satn.-binding expts. The pharmacol. profile using 125I-Cetorelix as radioligand and the authentic gonadoliberein or agonistic and antagonistic derivs. as competitors, showed a distinct rank order of binding potencies. Superagonistic gonadoliberein derivs. had more than 10 times higher binding affinities in comparison to gonadoliberein with a Kd of 3.47 nM. The gonadoliberein receptor expressed in stably transfected LTK- cells coupled to the **inositol** phosphate signal-transduction pathway. Gonadoliberein stimulated the synthesis of **inositol 1,4,5-trisphosphate** in a dose-dependent way with an EC50 of 5 nM. This stimulatory effect of gonadoliberein was completely antagonized by Cetorelix in equimolar concns., demonstrating the high potency of this competitive receptor antagonist. A transient expression of the c-fos protooncogene in growth-arrested cells was induced by gonadoliberein or [D-Trp6]gonadoliberein. The gonadoliberein receptor couples to a putative mitogenic signal-transduction pathway in this heterologous cell system.
ST gonadoliberein receptor cDNA cloning cell line; LHRH receptor characterization fibroblast cell line
IT Signal transduction, biological
(gonadoliberein receptors function and characterization after stable over-expression in fibroblast cell line)
IT Receptors
RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
(LH-releasing factor, gonadoliberein receptors function and characterization after stable over-expression in fibroblast cell line)
IT Animal cell line
(LTK-, gonadoliberein receptors function and characterization after stable over-expression in fibroblast cell line)
IT **Gene**, animal
RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
(c-fos, gonadoliberein receptors function and characterization after stable over-expression in fibroblast cell line)
IT 52435-06-0 57773-63-4 57982-77-1 112568-12-4, Antide 120287-85-6, Cetorelix 151272-78-5, Antarelix
RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
(gonadoliberein receptors function and characterization after stable over-expression in fibroblast cell line)
IT 9034-40-6, LH-RH
RL: BPR (Biological process); BSU (Biological study, unclassified); BIOL (Biological study); PROC (Process)
(gonadoliberein receptors function and characterization after stable over-expression in fibroblast cell line)
IT 88269-39-0, **Inositol 1,4,5-trisphosphate**
RL: BPR (Biological process); MFM (Metabolic formation); BIOL (Biological study); FORM (Formation, nonpreparative); PROC (Process)
(gonadoliberein receptors function and characterization after stable over-expression in fibroblast cell line)
L5 ANSWER 5 OF 8 CAPLUS COPYRIGHT 1999 ACS
AN 1995:358243 CAPLUS
DN 122:181931
TI src-Homology 2 (SH2) domain ligation as an allosteric regulator:

modulation of phosphoinositide-specific phospholipase C.gamma.1 structure and activity

AU Koblan, Kenneth S.; Schaber, Michael D.; Edwards, Gwynneth; Gibbs, Jackson B.; Pompliano, David L.

CS Dep. Cancer Res., Merck Res. Lab., West Point, PA, 19486, USA

SO Biochem. J. (1995), 305(3), 745-51
CODEN: BIJOAK; ISSN: 0264-6021

DT Journal

LA English

CC 7-5 (Enzymes)

Section cross-reference(s): 3

AB Phosphatidylinositol (PI)-specific phospholipase C.gamma.1 (PI-PLC.gamma.1) catalyzes the hydrolysis of PI 4,5-bisphosphate (PIP2) to generate the 2nd messengers diacylglycerol and **inositol 1,4,5-trisphosphate**. PI-PLC.gamma.1, an src-homol. 2/3 (SH2/SH3) domain-contg. enzyme, is activated in response to growth factor-induced tyrosine phosphorylation, and, in vivo, is translocated from the cytosol to the particulate cell fraction. Here, the authors report the bacterial (Escherichia coli) expression of rat brain PI-PLC.gamma.1 under the control of the phage T7 promoter. Prodn. of the active enzyme in amts. suitable for structure-function anal. depended on coupling the translation of PI-PLC.gamma.1 to the expression of the phage- ϕ 10 coat protein. Purifn. of the enzyme was facilitated by the presence of a 3-amino-acid C-terminal antibody epitope tag (Glu-Glu-Phe) engineered into the cloned PI-PLC.gamma.1. Examn. of the specific activity, pH-rate profile, [Ca²⁺]-dependence, and substrate specificity of bacterially expressed PI-PLC.gamma.1 indicated that it had kinetic properties similar to those of PI-PLC.gamma.1 isolated from bovine brain. The substrate specificity was dependent on [Ca²⁺]: at low [Ca²⁺] (1-10 μ M), PIP2 was a better substrate than PI. The addn. of phosphotyrosine-contg. peptides (12-mers) with the cognate sequence of the **high-affinity binding** site for PI-PLC.gamma.1 on the activated epidermal growth factor (EGF) receptor (Tyr-992) increased enzyme activity (up to 85%) in vitro. Cognate nonphosphorylated peptides had no effect on enzyme activity. When CD spectroscopy was used to monitor the effect of added phosphotyrosine-contg. peptide on the structure of recombinant PI-PLC.gamma.1, significant spectral shifts, indicative of a conformational change, were obsd. upon complexation with the EGF-receptor phosphotyrosine-contg. 12-residue peptide (Tyr*-992). How SH2 domains from PI-PLC.gamma.1 can mediate structural rearrangements and modulate enzymic activity on their ligation by growth factor receptors was discussed.

ST phosphoinositide phospholipase C recombinant form brain; conformation change recombinant phosphoinositide phospholipase C; Escherichia expression rat brain phospholipase C

IT Escherichia coli
(cloning and expression of rat brain phosphatidylinositol phospholipase C in Escherichia coli)

IT Conformation and Conformers
(modulation of recombinant rat brain phosphoinositide-specific phospholipase C.gamma.1 structure by phosphotyrosine-contg. phosphopeptides)

IT Brain
(src-homol. 2 (SH2) domain ligation as an allosteric regulator: modulation of recombinant rat brain phosphoinositide-specific phospholipase C.gamma.1 structure and activity)

IT Receptors
RL: BPR (Biological process); BIOL (Biological study); PROC (Process)
(epidermal growth factor/.alpha.-transforming growth factor,

gene c-erbB, src-homol. 2 (SH2) domain ligation as an allosteric regulator: modulation of recombinant rat brain phosphoinositide-specific phospholipase C.gamma.1 structure and activity)

IT Phosphopeptides
 RL: BPR (Biological process); BIOL (Biological study); PROC (Process) (phosphotyrosine-contg., modulation of recombinant rat brain phosphoinositide-specific phospholipase C.gamma.1 structure by phosphotyrosine-contg. phosphopeptides)

IT Animal growth regulator receptors
 RL: BPR (Biological process); BIOL (Biological study); PROC (Process) (.alpha.-transforming growth factor **gene** c-erbB, src-homol. 2 (SH2) domain ligation as an allosteric regulator: modulation of recombinant rat brain phosphoinositide-specific phospholipase C.gamma.1 structure and activity)

IT 63551-76-8P, Phosphatidylinositol phospholipase C
 RL: BPN (Biosynthetic preparation); PEP (Physical, engineering or chemical process); PRP (Properties); PUR (Purification or recovery); BIOL (Biological study); PREP (Preparation); PROC (Process) (isoform .gamma.1; src-homol. 2 (SH2) domain ligation as an allosteric regulator: modulation of recombinant rat brain phosphoinositide-specific phospholipase C.gamma.1 structure and activity)

L5 ANSWER 6 OF 8 CAPLUS COPYRIGHT 1999 ACS
 AN 1993:18143 CAPLUS
 DN 118:18143
 TI Structure of a novel InsP3 receptor
 AU Sudhof, Thomas C.; Newton, Christopher L.; Archer, Branch T., III; Ushkaryov, Yuri A.; Mignery, Gregory A.
 CS Southwest. Med. Cent., Univ. Texas, Dallas, TX, 75235, USA
 SO EMBO J. (1991), 10(11), 3199-206
 CODEN: EMJODG; ISSN: 0261-4189
 DT Journal
 LA English
 CC 6-3 (General Biochemistry)
 Section cross-reference(s): 2, 3, 13

AB **Inositol 1,4,5-trisphosphate** (InsP3) constitutes a major intracellular second messenger that transduces many growth factor and neurotransmitter signals.
 InsP3 causes the release of Ca²⁺ from intracellular stores by binding to specific receptors that are coupled to Ca²⁺ channels. One such receptor from cerebellum has previously been extensively characterized. The authors have now detd. the full structure of a second, novel InsP3 receptor which is referred to as type 2 InsP3 receptor as opposed to the cerebellar type 1 InsP3 receptor. The type 2 InsP3 receptor has the same general structural design as the cerebellar type 1 InsP3 receptor with which it shares 69% sequence identity. Expression of the N-terminal 1078 amino acids of the type 2 receptor demonstrates **high-affinity binding** of InsP3 to the type 2 receptor with a similar specificity but higher affinity than obsd. for the type 1 receptor. These results demonstrate the presence of several types of InsP3 receptor in brain and raise the possibility that intracellular Ca²⁺ signaling may involve multiple pathways with different regulatory properties dependent on different InsP3 receptors.

ST rat **inositol** trisphosphate receptor **gene**;
inositol trisphosphate receptor sequence cerebellum; protein
inositol trisphosphate binding cerebellum

IT **Gene**, animal
 RL: BIOL (Biological study)
 (for **inositol** tris(phosphate)receptor type 2, of rat cerebellum, nucleotide sequence of)

IT Rat
 (**inositol** tris(phosphate) receptor type 2 **gene** of

cerebellum of, nucleotide sequence of)
 IT Deoxyribonucleic acid sequences
 (**inositol** tris(phosphate)-binding protein-specifying, type 2,
 of cerebellum, of rat, complete)
 IT Protein sequences
 (of **inositol** tris(phosphate) receptor type II, of rat
 cerebellum, complete)
 IT Brain, composition
 (cerebellum, **inositol** tris(phosphate) receptor type 2 of,
 amino acid sequence of)
 IT Receptors
 RL: BIOL (Biological study)
 (**inositol** tris(phosphate), type 2, amino acid sequence of, of
 cerebellum)
 IT Proteins, specific or class
 RL: BIOL (Biological study)
 (**inositol** tris(phosphate)-binding, type 2, amino acid
 sequence of, of cerebellum)
 IT 145112-31-8, Protein (rat clone p547-13/pI71/pI70/pI53/pI15/pI6
 inositol tris(phosphate)-binding isoform 2 reduced)
 RL: PRP (Properties)
 (amino acid sequence of)
 IT 140991-42-0 145112-30-7
 RL: PRP (Properties)
 (nucleotide sequence of)
 IT 88269-39-0, **Inositol 1,4,5-**
 trisphosphate
 RL: BIOL (Biological study)
 (receptor for, type 2, amino acid sequence of, of cerebellum of rat)

L5 ANSWER 7 OF 8 CAPLUS COPYRIGHT 1999 ACS

AN 1992:100352 CAPLUS

DN 116:100352

TI Cloning and expression of a complementary DNA encoding a bovine adrenal
 angiotensin II type-1 receptor

AU Sasaki, Katsutoshi; Yamano, Yoshiaki; Bardhan, Smriti; Iwai, Naoharu;
 Murray, John J.; Hasegawa, Mamoru; Matsuda, Yuzuru; Inagami, Tadashi

CS Dep. Biochem., Vanderbilt Univ., Nashville, TN, 37232, USA

SO Nature (London) (1991), 351(6323), 230-3

CODEN: NATUAS; ISSN: 0028-0836

DT Journal

LA English

CC 3-3 (Biochemical Genetics)

Section cross-reference(s): 6, 13

AB The expression cloning of a cDNA encoding a bovine angiotensin II
 receptor

is reported. The receptor cDNA encodes a protein of 359 amino-acid
 residues with a transmembrane topol. similar to that of other G
 protein-coupled receptors. COS-7 cells transfected with the cDNA
 expressed specific and **high-affinity binding**
 sites for angiotensin II, angiotensin II antagonist and a non-peptide
 specific antagonist for type-1 receptor. Dithiothreitol inhibited ligand
 binding. The concn. of intracellular Ca²⁺ and of **inositol-**
1,4,5-trisphosphate increased in the
 transfected COS-7 cells in response to angiotensin II or angiotensin III,
 indicating that this receptor is the type-1 receptor for angiotensin II.
 Northern blot anal. revealed that the mRNA for this receptor is expressed
 in bovine adrenal medulla, cortex and kidney.

ST cattle angiotensin receptor cDNA sequence

IT Cattle

(angiotensin II type-1 receptor **gene** of, cloning and
 nucleotide and encoded peptide sequences of)

IT Adrenal gland, composition

(angiotensin II type-1 receptor **gene** of, of cattle, cloning
 and nucleotide and encoded peptide sequences of)

IT **Gene**, animal

RL: BIOL (Biological study)
 (for angiotensin II type-1 receptor, of cattle, cloning and nucleotide
 and encoded peptide sequences of)
 IT Molecular cloning
 (of angiotensin II type-1 receptor **gene**, of cattle)
 IT Protein sequences
 (of angiotensin II type-1 receptor, of cattle, complete)
 IT Deoxyribonucleic acid sequences
 (receptor-specifying, type 1, for angiotensin II, of cattle, complete)
 IT Receptors
 RL: BIOL (Biological study)
 (type-1, for angiotensin II, **gene** for, of cattle, cloning and
 nucleotide and encoded peptide sequences of)
 IT Animal cell line
 (COS-7, cattle angiotensin II type-1 receptor **gene** expression
 in)
 IT 7440-70-2, Calcium, biological studies
 RL: BIOL (Biological study)
 (COS-7 cell increase of, in angiotensin presence, after cattle
 angiotensin II type-1 receptor **gene** cloning)
 IT 88269-39-0, **Inositol**-1,4,5-triphosphate
 RL: PROC (Process)
 (COS-7 cell increase of, in angiotensin presence, after cattle
 angiotensin II type-1 receptor **gene** cloning)
 IT 138464-42-3
 RL: PRP (Properties)
 (amino acid sequence of)
 IT 138440-38-7, Deoxyribonucleic acid (ox clone ARW angiotensin II receptor
 isoform AT1 messenger RNA-complementary) 138440-39-8
 RL: BIOL (Biological study); PRP (Properties)
 (nucleotide sequence of)
 IT 11128-99-7, Angiotensin II
 RL: PRP (Properties)
 (receptor type-1 for, **gene** for, of cattle, cloning and
 nucleotide and encoded peptide sequences of)
 IT 9041-90-1, Angiotensin I 9088-01-1 11130-03-3, [Sar1,
 Ala8]-Angiotensin II 12687-51-3, Angiotensin III 124750-99-8, Dup753
 RL: PRP (Properties)
 (recombinant bovine angiotensin II type-1 receptor binding by)

L5 ANSWER 8 OF 8 CAPLUS COPYRIGHT 1999 ACS

AN 1989:627531 CAPLUS

DN 111:227531

TI Four intracisternal calcium-binding glycoproteins from rat liver
 microsomes with high affinity for calcium. No indication for
 calsequestrin-like proteins in **inositol 1,4,**
5-trisphosphate-sensitive calcium sequestering rat liver
 vesicles

AU Nguyen Van Phuc; Peter, Frank; Soeling, Hans Dieter

CS Zent. Inn. Med., Univ. Goettingen, Goettingen, D-3400, Fed. Rep. Ger.

SO J. Biol. Chem. (1989), 264(29), 17494-501

CODEN: JBCHA3; ISSN: 0021-9258

DT Journal

LA English

CC 6-3 (General Biochemistry)

AB The **inositol 1,4,5-**

trisphosphate-sensitive compartment of rat liver does not contain
 calsequestrin-like material. Instead 4 nonmembranous Ca²⁺-binding
 glycoproteins with approx. mol. masses of 59, 60, 80, and 90 kDa were
 found. The 59-, 80-, and 90-kDa proteins were of the high mannose-rich
 type, the carbohydrate moiety of the 60-kDa protein was of the complex
 hybrid type with terminal galactoses. All 4 proteins had **high**
affinity binding sites for Ca²⁺ (KD 1-5 .mu.M) and 1-5
 binding sites/mol. The 80- and the 90-kDa proteins also had low affinity
 binding sites (KD 400 and 600 .mu.M, resp., with 13 and 15 binding
 sites/mol., resp.). A comparison of the N-terminal sequences revealed

that the 60-kDa Ca²⁺-binding protein represents the rat liver calregulin, whereas the 90-kDa Ca²⁺-binding protein represents grp94. The sequences did not reveal any relationship of the 80-kDa protein with grp78, or of the 59-kDa protein with protein disulfide isomerase.

ST calcium binding glycoprotein liver; calregulin calcium binding liver;
gene grp94 glycoprotein calcium binding liver

IT Liver, composition
 (calcium-binding glycoproteins of endoplasmic reticulum of)

IT Calsequestrins
 RL: BIOL (Biological study)
 (calcium-binding glycoproteins of liver endoplasmic reticulum in relation to)

IT Endoplasmic reticulum
 (calcium-binding glycoproteins of, of liver)

IT Protein sequences
 (of calregulin N terminus, of liver)

IT Conformation and Conformers
 (of calregulin and grp78 glycoprotein, of liver endoplasmic reticulum)

IT Oligosaccharides
 RL: BIOL (Biological study)
 (of calregulin and grp94 glycoprotein, of liver endoplasmic reticulum)

IT Protein sequences
 (of grp94 glycoprotein N terminus, of liver)

IT Glycoproteins, specific or class
 RL: BIOL (Biological study)
 (calcium-binding, 59,000-mol.-wt., of endoplasmic reticulum, of liver)

IT Glycoproteins, specific or class
 RL: BIOL (Biological study)
 (calcium-binding, 80,000-mol.-wt., of endoplasmic reticulum, of liver)

IT Glycoproteins, specific or class
 RL: BIOL (Biological study)
 (calreticulins, of endoplasmic reticulum, of liver, carbohydrate structure of and other calcium-binding glycoproteins comparison with)

IT Glycoproteins, specific or class
 RL: BIOL (Biological study)
 (endoplasmins, of endoplasmic reticulum, of liver, carbohydrate structure of and other calcium-binding glycoproteins comparison with)

IT 88269-39-0, **Inositol** 1,4,5-triphosphate
 RL: BIOL (Biological study)
 (calcium release from liver endoplasmic reticulum sensitive to, calcium-binding glycoproteins in relation to)

IT 7440-70-2, Calcium, biological studies
 RL: BIOL (Biological study)

L17 ANSWER 1 OF 2 MEDLINE
 AN 97303864 MEDLINE
 DN 97303864
 TI Binding and activity of the nine possible regioisomers of myo-
inositol tetrakisphosphate at the **inositol 1, 4, 5-trisphosphate** receptor.
 AU Burford N T; Nahorski S R; Chung S K; Chang Y T; Wilcox R A
 CS Department of Cell Physiology and Pharmacology, University of Leicester, UK.
 SO CELL CALCIUM, (1997 Apr) 21 (4) 301-10.
 Journal code: CQE. ISSN: 0143-4160.
 CY SCOTLAND: United Kingdom
 DT Journal; Article; (JOURNAL ARTICLE)
 LA English
 FS Priority Journals
 EM 199710
 EW 19971001
 AB All 9 racemic regioisomers (15 enantiomerically) of myo-**inositol** tetrakisphosphates (IP4s): DL-Ins(1,2,4,5)P4 [A], DL-Ins(1,2,4,6)P4 [B], Ins(1,2,3,5)P4 [C], Ins(1,3,4,6)P4 [D], Ins(2,4,5,6)P4 [E], DL-Ins(1,3,4,5)P4 [F], DL-Ins(1,2,5,6)P4 [G], DL-Ins(1,2,3,4)P4 [H] and DL-Ins(1,4,5,6)P4 [I] [Chung S-K., Chang Y-T. Synthesis of all possible regioisomers of myo-**inositol** tetrakisphosphate. J Chem Soc Chem Commun 1995; 11-13] were investigated for their ability to bind to the D-myo-**inositol 1, 4, 5-trisphosphate** [Ins(1,4,5)P3] receptor in bovine adrenal cortical membranes, and for their ability to mobilize 45Ca2+ from Ins(1,4,5)P3-sensitive Ca2+ stores in permeabilized Chinese hamster ovary (CHO) cells. DL-Ins(1,2,4,5)P4 (Ki = 11 nM) bound to Ins(1,4,5)P3 receptors with an affinity only 2-fold lower than Ins(1,4,5)P3 (Ki = 6 nM). Ins(1,2,3,5)P4, Ins(1,3,4,6)P4, Ins(2,4,5,6)P4, DL-Ins(1,3,4,5)P4, DL-Ins(1,2,3,4)P4 and DL-Ins(1,4,5,6)P4 bound with affinities of between 0.4-0.7 microM. DL-Ins(1,2,4,6)P4 and DL-Ins(1,2,5,6)P4 bound to the Ins(1,4,5)P3 receptor with low affinity (approximately 2-3 microM). All but one of the IP4s mediated release of 45Ca2+ from stores of permeabilized CHO cells with a similar rank order of potency as that for Ins(1,4,5)P3 receptor binding, being between 16-fold and 50-fold less potent at releasing 45Ca2+ compared with their apparent binding affinities to the Ins(1,4,5)P3 receptor. The notable exception was Ins(1,2,3,5)P4, which showed an approximately 200-fold lower potency compared with its affinity for the Ins(1,4,5)P3 receptor. Ins(1,2,3,5)P4 may be a useful lead compound for the rational design of novel synthetic Ins(1,4,5)P3 analogues possessing structure-activity profiles with relatively **high binding** affinity, but low intrinsic efficacy, and hence partial agonists and antagonists at the Ins(1,4,5)P3 receptor.
 CT Check Tags: Animal; Support, Non-U.S. Gov't
 Calcium: ME, metabolism
 *Calcium Channels: ME, metabolism
 Cattle
 Cell Membrane Permeability
 CHO Cells
 Hamsters
 *Inositol Phosphates: ME, metabolism
 Ionomycin: PD, pharmacology
 Ionophores: PD, pharmacology
 Isomerism
 Models, Chemical
 *Receptors, Cytoplasmic and Nuclear: ME, metabolism
 RN 102850-29-3 (**inositol-1,3,4,5-tetrakisphosphate**); 56092-81-0

(Ionomycin); 7440-70-2 (Calcium)
 CN 0 (**inositol**-1,4,5-triphosphate receptor); 0 (Calcium Channels);
 0 (**Inositol** Phosphates); 0 (Ionophores); 0 (Receptors,
 Cytoplasmic and Nuclear)

L17 ANSWER 2 OF 2 MEDLINE
 AN 96074809 MEDLINE
 DN 96074809
 TI Molecular cloning and expression of multiple isoforms of human
 prostaglandin E receptor EP3 subtype generated by alternative messenger
 RNA splicing: multiple second messenger systems and tissue-specific
 distributions.
 AU Kotani M; Tanaka I; Ogawa Y; Usui T; Mori K; Ichikawa A; Narumiya S;
 Yoshimi T; Nakao K
 CS Department of Medicine, Faculty of Medicine, Kyoto University, Japan.
 SO MOLECULAR PHARMACOLOGY, (1995 Nov) 48 (5) 869-79.
 Journal code: NGR. ISSN: 0026-895X.
 CY United States
 DT Journal; Article; (JOURNAL ARTICLE)
 LA English
 FS Priority Journals; Cancer Journals
 OS GENBANK-D38297; GENBANK-D38298; GENBANK-D38299; GENBANK-D38300;
 GENBANK-D38301
 EM 199602
 AB Five distinct cDNA clones encoding four different isoforms of human
 prostaglandin (PG) E receptor EP3 subtype were isolated from a human
 kidney cDNA library. Two cDNA clones differed only in their
 3'-untranslated regions. The four isoforms, tentatively named EP3-I,
 EP3-II, EP3-III, and EP3-IV, which were generated by alternative mRNA
 splicing, had identical amino acid sequences except for their different
 carboxyl-terminal tails. Transfection experiments revealed that all the
 four isoforms show **high binding** affinities to PGE2,
 PGE1, and M&B28767, an EP3-specific agonist, whereas their downstream
 signaling pathways are divergent. M&B28767 increased cAMP concentrations
 in cells expressing EP3-II and EP3-IV, whereas it inhibited
 forskolin-induced cAMP accumulations in cells expressing all EP3
 isoforms.
 M&B28767 also stimulated phosphoinositide turnover in cells expressing
 EP3-I and EP3-II. Northern blot analysis revealed that the EP3 gene is
 expressed in a wide variety of human tissues. The human EP3 mRNA was
 present most abundantly in the kidney, pancreas, and uterus. A
 substantial
 expression was also detected in the heart, liver, skeletal muscle, small
 intestine, colon, prostate, ovary, and testis. Furthermore, reverse
 transcription-polymerase chain reaction analysis demonstrated
 tissue-specific expressions of the five different EP3 mRNA species. The
 present study suggests the presence of the multiple systems of PGE2/EP3
 isoforms and leads to the better understanding of its physiological and
 pathophysiological implications in humans.
 CT Check Tags: Animal; Human; Support, Non-U.S. Gov't
 *Alternative Splicing
 Amino Acid Sequence
 Base Sequence
 Blotting, Northern
 Blotting, Southern
 Cloning, Molecular
 Cyclic AMP: AN, analysis
 CHO Cells
 Dinoprostone: ME, metabolism
 Hamsters
Inositol 1,4,5-Trisphosphate: AN, analysis
 Molecular Sequence Data
 Organ Specificity
 Polymerase Chain Reaction
 Receptors, Prostaglandin E: CH, chemistry
 *Receptors, Prostaglandin E: GE, genetics

Receptors, Prostaglandin E: PH, physiology
*Second Messenger Systems
*Second Messenger Systems: PH, physiology
RN 363-24-6 (Dinoprostone); 60-92-4 (Cyclic AMP); **85166-31-0 (Inositol
1,4,5-Trisphosphate)**
CN 0 (Receptors, Prostaglandin E)

29 ANSWER 1 OF 3 MEDLINE
 AN 1999119984 MEDLINE
 DN 99119984
 TI Interaction of the Na(+)-Ca2+ exchanger with small molecules on cell Ca2+ signaling.
 AU Fang Y; Rong M; He L
 CS Department of Anesthesiology, Zhong Shan Hospital, China.
 SO BIOMEDICINE AND PHARMACOTHERAPY, (1998) 52 (10) 459-64.
 Journal code: A59. ISSN: 0753-3322.
 CY France
 DT Journal; Article; (JOURNAL ARTICLE)
 LA English
 FS Priority Journals
 EM 199906
 EW 19990603
 AB Interactions of the Na(+)-Ca2+ exchanger with small molecules on cell Ca2+ signaling were elucidated in Chinese hamster ovary (CHO) C1 cells, which transfected a control **vector** without any expression of the Na(+)-Ca2+ exchanger's gene while CHO CK1.4 cells transfected an expression **vector** encoding the bovine cardiac Na(+)-Ca2+ exchanger's cDNA, treated with lithium- or sodium-buffer medium respectively, by using L16(2)15 multifactorial orthogonal statistics and fura-2 fluorescence real-time imaging. In contrast to controls of Li(+)-treated C1 cells, the store-dependent Ca(2+)-influx (SDCI) was enhanced by either the Na(+)-Ca2+ exchanger, Na(+), 1-((beta-[3-(4-methoxyphenyl)propoxy]-4-methoxyphenethyl)-1H-imidazole HCl (SK&F96365) or ouabain, and by interactions of the Na(+)-Ca2+ exchanger with either Na+, SK&F96365 or both SK&F96365 and ouabain; and ATP-induced Ca2+ release (AICR) was activated by SK&F96365 or Na+ alone, interactions of the Na(+)-Ca2+ exchanger with SK&F96365 or Na+, and an interaction between SK&F96365 and ouabain. The dramatic interaction of the Na(+)-Ca2+ exchanger with small molecules indicates that cell Ca2+ signaling is generated by **inositol** triphosphate (InsP3)-dependent pathways, allosteric effects of the G-protein coupled P2y&2u purinoceptor and multi-site recognition. Our findings provide meaningful clues for designing new strategies of cardiocerebral vascular oxidative **diseases**.
 CT Check Tags: Animal; Support, Non-U.S. Gov't
 Adenosine Triphosphate: PD, pharmacology
 *Calcium Signaling: PH, physiology
 CHO Cells
 Fluorescent Dyes
 Fura-2
 Hamsters
 Indicators and Reagents
 *Sodium-Calcium Exchanger: PH, physiology
 RN 56-65-5 (Adenosine Triphosphate); 96314-98-6 (Fura-2)
 CN 0 (Fluorescent Dyes); 0 (Indicators and Reagents); 0 (Sodium-Calcium Exchanger)

 L29 ANSWER 2 OF 3 MEDLINE
 AN 94253977 MEDLINE
 DN 94253977
 TI Correlation of phosphoinositide hydrolysis with exflagellation in the malaria microgametocyte.
 AU Martin S K; Jett M; Schneider I
 CS Department of Molecular Pathology, Walter Reed Army Institute of Research,

Washington, DC 20307-5100.

SO JOURNAL OF PARASITOLOGY, (1994 Jun) 80 (3) 371-8.
Journal code: JL3. ISSN: 0022-3395.

CY United States

DT Journal; Article; (JOURNAL ARTICLE)

LA English

FS Priority Journals

EM 199409

AB Cellular responses to growth factors, hormones, and other agonists have been shown in many animal cell systems to be mediated by the signal transduction cascade controlled by phospholipase C. One such response, calcium mobilization, is regulated by the concerted effect of several specific **inositol** (poly)phosphates. Another response, protein phosphorylation, is regulated by other phospholipase C (PLC) hydrolysis products. Mature gametocytes are specialized cells primed for transformation into gametes immediately upon removal from the vertebrate bloodstream, thereby initiating the sexual cycle in a **vector** mosquito. This study showed that PLC hydrolysis products, **inositol** (1,4,5)triphosphate and diacylglycerol, are correlated with the initial events of flagellar development; they are implicated in synchronizing this crucial transformation for the parasite and hence the continued transmission of the parasite, which leads to this debilitating **disease**.

CT Check Tags: Animal
Chromatography, High Pressure Liquid
Culture Media
Diglycerides: ME, metabolism
Diglycerides: PH, physiology
*Flagella: PH, physiology
Hydrolysis
Inositol 1,4,5-Trisphosphate: ME, metabolism
Inositol 1,4,5-Trisphosphate: PH, physiology
Kinetics
Microscopy, Phase-Contrast
*Phosphatidylinositols: ME, metabolism
Phospholipase C: PH, physiology
Plasmodium falciparum: ME, metabolism
*Plasmodium falciparum: PH, physiology
Plasmodium falciparum: UL, ultrastructure
Signal Transduction
Substrate Specificity

RN **85166-31-0 (Inositol 1,4,5-Trisphosphate)**

CN EC 3.1.4.3 (Phospholipase C); 0 (Culture Media); 0 (Diglycerides); 0 (Phosphatidylinositols)

L29 ANSWER 3 OF 3 MEDLINE

AN 94018073 MEDLINE

DN 94018073

TI Multiple regression of skeletal muscle tension on **inositol** phosphates: cross-talk between signal transduction mechanisms in burn trauma.

AU Tomera J F; Lilford K

CS Clinical Pharmacology Laboratory, Shriners Burn Institute, Boston, MA.

SO METHODS AND FINDINGS IN EXPERIMENTAL AND CLINICAL PHARMACOLOGY, (1993 Jun)
15 (5) 255-65.
Journal code: LZN. ISSN: 0379-0355.

CY Spain

DT Journal; Article; (JOURNAL ARTICLE)

LA English

FS Priority Journals

EM 199401

AB Skeletal muscle weakness associated with burn trauma prolongs the time of rehabilitation of burn patients. Understanding the underlying chemical changes that impact on physiological tension may provide new therapeutic

options for the treatment of burn patients. This report demonstrates the novelty of applying 3-dimensional graphic capabilities, involving area and **vector** changes to understand variations in **inositol** derivatives and their co-modulating influence on physiological tension in skeletal muscle. This muscle was distant from the primary anatomical burn site. It was subjected to circulatory shock emanating from burn trauma. Burn injury was achieved by scalding of predefined areas (0, 20% and 50%) on the dorsal and ventral surfaces of mice. At day 21, tension studies via muscle twitch analyses were performed. Through multiple regression, the dependency of physiologic tension was determined with respect to three poly-**inositol** forms each representing independent parameters simultaneously. The contribution of each of these parameters was assigned to a three-dimensional axis. Relationships of tension on three fixed independent parameters were found only for the 20% and 50% burn groups. **Vector** analysis on a plane in three-dimensional space determined the relationship of tension to each of the independent parameters in 20% and 50% burn groups. No significant relationship of tension dependency on three fixed poly-**inositol** variables was found in the control group. Such **vector** analysis, using solid and differential analytical geometry, allowed for a clear visualization of the interrelationships that existed between secondary messenger systems (viz, IP3) and a resulting physiologic manifestation (viz, tension). This clear that visualization allows for a greater understanding of messenger systems that may lead to more effective treatment of skeletal muscle weakness associated with the systemic effects of severe burn trauma.

CT Check Tags: Animal; Male; Support, Non-U.S. Gov't
 Burns: ME, metabolism
 *Burns: PP, physiopathology
 Disease Models, Animal
 Inositol Phosphates: ME, metabolism
 *Inositol Phosphates: PH, physiology
 Inositol 1,4,5-Trisphosphate: ME, metabolism
 Inositol 1,4,5-Trisphosphate: PD, pharmacology
 Mice
 Mice, Inbred Strains
 Multivariate Analysis
 Muscle Contraction: DE, drug effects
 Muscle Contraction: PH, physiology
 Muscles: DE, drug effects
 Muscles: ME, metabolism
 *Muscles: PP, physiopathology
 Regression Analysis
 *Signal Transduction: PH, physiology

RN 85166-31-0 (Inositol 1,4,5-Trisphosphate)
 CN 0 (Inositol Phosphates)